SEQUENCE LISTING

<110> Human Genome Sciences, Inc.

<120> Keratinocyte Derived Interferon

<130> PF482P1

T2540

<140> Unassigned
<141> 2000-01-20

<150> 60/093,643
<151> 1998-07-21

<150> PCT/US99/16424
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<170> PatentIn Ver. 2.1

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50 55 60

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Ile Phe Ser Gln His Thr Phe Lys Tyr Trp Lys Glu Arg His Leu Lys
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35 40 45

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65 70 75 80

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85 90 95

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100 105 110

His Gln Gln Leu Gln His Leu Glu Thr Cys Leu Leu Gln Val Val Gly
115 120 125

Glu Gly Glu Ser Ala Gly Ala Ile Ser Ser Val Pro Gln Leu Ser Ser
130 135 140

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145 150 155 160

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165 170 175

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180 185 190

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195 200 205

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Leu Glu Tyr Cys Leu Lys Asp Arg Met Asn Phe Asp Ile Pro Glu Glu
50 55 60

Ile Lys Gln Leu Gln Gln Phe Gln Lys Glu Asp Ala Ala Leu Thr Ile
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Tyr Glu Met Leu Gln Asn Ile Phe Ala Ile Phe Arg Gln Asp Ser Ser
85 90 95

Ser Thr Gly Trp Asn Glu Thr Ile Val Glu Asn Leu Leu Ala Asn Val
100 105 110

Tyr His Gln Ile Asn His Leu Lys Thr Val Leu Glu Glu Lys Leu Glu
115 120 125

Lys Glu Asp Phe Thr Arg Gly Lys Leu Met Ser Ser Leu His Leu Lys
130 135 140

Arg Tyr Tyr Gly Arg Ile Leu His Tyr Leu Lys Ala Lys Glu Tyr Ser
145 150 155 160

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35 40 45

Pro His Phe Cys Leu Gln Asp Arg Lys Asp Phe Ala Leu Pro Gln Glu
50 55 60

Met Val Glu Gly Gly Gln Leu Gln Glu Ala Gln Ala Ile Ser Val Leu
65 70 75 80

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85 90 95

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100 105 110

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115 120 125

Glu Glu Asp Ser Ala Leu Gly Arg Thr Gly Pro Leu Ala Leu Lys Arg
130 135 140

Tyr Phe Gln Gly Ile His Val Tyr Leu Lys Glu Lys Gly Tyr Ser Asp
145 150 155 160

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180 185 190

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35 40 45

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50 55 60

Met Val Lys Gly Ser Gln Leu Gln Lys Ala His Val Met Ser Val Leu
65 70 75 80

His Glu Met Leu Gln Gln Ile Phe Ser Leu Phe His Thr Glu Arg Ser
85 90 95

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100 105 110

His Gln Gln Leu Gln His Leu Glu Thr Cys Leu Leu Gln Val Val Gly
115 120 125

Glu Gly Glu Ser Ala Gly Ala Ile Ser Ser Val Pro Gln Leu Ser Ser
130 135 140

Leu Glu Leu Arg Arg Tyr Phe His Arg Ile Asp Asn Phe Leu Lys Glu
145 150 155 160

Lys Lys Tyr Ser Asp Cys Ala Trp Glu Ile Val Arg Val Glu Ile Arg
165 170 175

Arg Cys Leu Tyr Tyr Phe Tyr Lys Phe Thr Ala Leu Pro Ala Leu Thr
180 185 190

Leu Arg Arg Tyr Phe Gln Gly Ile Arg Val Tyr Leu Lys Glu Lys Lys
195 200 205

Tyr Ser Asp Cys Ala Trp Glu Val Val Arg Met Glu Ile Met Lys Ser
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Asp Leu Gly Ser Ser
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35 40 45

Pro Phe Ser Cys Leu Lys Asp Arg His Asp Phe Arg Ile Pro Gln Glu
50 55 60

Glu Phe Asp Gly Asn Gln Phe Gln Asp Ala Gln Ala Ile Ser Val Leu
65 70 75 80

His Glu Met Ile Gln Gln Thr Phe Asn Leu Phe Ser Thr Glu Asp Ser
85 90 95

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Tyr Gln Gln Leu Asn Asp Leu Glu Ala Cys Val Ile Gln Glu Val Gly
        115                 120                 125

Val Glu Glu Thr Pro Leu Met Asn Glu Asp Ser Ile Leu Ala Val Arg
    130                 135                 140

Lys Tyr Phe Gln Arg Ile Thr Leu Tyr Leu Ile Glu Arg Lys Tyr Ser
145                 150                 155                 160

Pro Cys Ala Trp Glu Val Val Arg Ala Glu Ile Met Arg Ser Leu Ser
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Phe Ser Thr Asn Leu Gln Lys Arg Leu Arg Arg Lys Asp
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Gly Asn Arg Arg Ala Leu Ile Leu Leu Ala Gln Met Gly Arg Ile Ser
        35                  40                  45

Pro Phe Ser Cys Leu Lys Asp Arg His Asp Phe Gly Phe Pro Gln Glu
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Glu Phe Asp Gly Asn Gln Phe Gln Lys Ala His Val Met Ser Val Leu
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His Glu Met Leu Gln Gln Ile Phe Ser Leu Phe His Thr Glu Arg Ser
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Ser Ala Ala Trp Glu Gln Ser Leu Leu Glu Lys Phe Ser Thr Glu Leu
            100                 105                 110

Asn Gln Gln Leu Asn Asp Leu Glu Ala Cys Val Ile Gln Glu Val Gly
        115                 120                 125

Val Glu Glu Thr Pro Leu Met Asn Val Asp Ser Ile Leu Ala Val Lys
    130                 135                 140

Lys Tyr Phe Gln Arg Ile Thr Leu Tyr Leu Thr Glu Lys Lys Tyr Ser
145                 150                 155                 160

Pro Cys Ala Trp Glu Val Val Arg Ala Glu Ile Met Arg Ser Phe Ser
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Leu Ser Lys Ile Phe Gln Glu Arg Leu Arg Arg Lys Glu
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35 40 45

Pro Phe Leu Cys Leu Lys Asp Arg Arg Asp Phe Arg Phe Pro Gln Glu
50 55 60

Met Val Lys Gly Ser Gln Leu Gln Lys Ala His Val Met Ser Val Leu
65 70 75 80

His Glu Met Leu Gln Gln Ile Phe Ser Leu Phe His Thr Glu Arg Ser
85 90 95

Ser Ala Ala Trp Asn Met Thr Leu Leu Asp Gln Leu His Thr Glu Leu
100 105 110

His Gln Gln Leu Gln His Leu Glu Thr Cys Leu Leu Gln Val Val Gly
115 120 125

Glu Gly Glu Ser Ala Gly Ala Ile Ser Ser Pro Ala Leu Thr Leu Arg
130 135 140

Arg Tyr Phe Gln Gly Ile Arg Val Tyr Leu Lys Glu Lys Lys Tyr Ser
145 150 155 160

Asp Cys Ala Trp Glu Val Val Arg Met Glu Ile Met Lys Ser Leu Phe
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180 185 190

Gly Ser Ser
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<213> Homo sapiens

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Gly Asn Arg Arg Ala Trp Ile Leu Leu Ala Gln Met Gly Arg Ile Ser
35 40 45

His Phe Ser Cys Leu Lys Asp Arg Tyr Asp Phe Gly Phe Pro Gln Glu
50 55 60

Val Phe Asp Gly Asn Gln Phe Gln Lys Ala Gln Ala Ile Ser Ala Phe
65 70 75 80

His Glu Met Ile Gln Gln Thr Phe Asn Leu Phe Ser Thr Lys Asp Ser
85 90 95

Ser Ala Ala Trp Asp Glu Thr Leu Leu Asp Lys Phe Tyr Ile Glu Leu
100 105 110

Phe Gln Gln Leu Asn Asp Leu Glu Ala Cys Val Thr Gln Glu Val Gly
115 120 125

Val Glu Glu Ile Ala Leu Met Asn Glu Asp Ser Ile Leu Ala Val Arg
130 135 140

Lys Tyr Phe Gln Arg Ile Thr Leu Tyr Leu Met Gly Lys Lys Tyr Ser
145 150 155 160

Pro Cys Ala Trp Glu Val Val Arg Ala Glu Ile Met Arg Ser Phe Ser
165 170 175

Phe Ser Thr Asn Leu Gln Lys Gly Leu Arg Arg Lys Asp
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Gly Ala Arg Glu Asn Leu Arg Leu Leu Ala Arg Met Asn Arg Leu Ser
35 40 45

Pro His Pro Cys Leu Gln Asp Arg Lys Asp Phe Gly Leu Pro Gln Glu
50 55 60

Met Val Glu Gly Asn Gln Leu Gln Lys Asp Gln Ala Ile Ser Val Leu
65 70 75 80

His Glu Met Leu Gln Gln Cys Phe Asn Leu Phe Tyr Thr Glu His Ser
85 90 95

Ser Ala Ala Trp Asn Thr Thr Leu Leu Glu Gln Leu Cys Thr Gly Leu
100 105 110

Gln Gln Gln Leu Glu Asp Leu Asp Ala Cys Leu Gly Pro Val Met Gly
115 120 125

Glu Lys Asp Ser Asp Met Gly Arg Met Gly Pro Ile Leu Thr Val Lys
130 135 140

Lys Tyr Phe Gln Gly Ile His Val Tyr Leu Lys Glu Lys Glu Tyr Ser

262

145 150 155 160

Asp Cys Ala Trp Glu Ile Ile Arg Met Glu Met Met Arg Ala Leu Ser
165 170 175

Ser Ser Thr Thr Leu Gln Lys Arg Leu Arg Lys Met Gly Gly Asp Leu
180 185 190

Asn Ser Leu
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20 25 30

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35 40 45

Pro His Ser Cys Leu Gln Asp Arg Lys Asp Phe Gly Leu Pro Gln Glu
50 55 60

Met Val Glu Gly Asp Gln Leu Gln Lys Asp Gln Ala Phe Pro Val Leu
65 70 75 80

Tyr Glu Met Leu Gln Gln Thr Phe Asn Leu Phe His Thr Glu His Ser
85 90 95

Ser Ala Ala Trp Asp Thr Thr Leu Leu Glu Gln Leu Cys Thr Gly Leu
100 105 110

Gln Gln Gln Leu Glu Asp Leu Asp Thr Cys Cys Arg Gly Gln Val Met
115 120 125

Gly Glu Glu Asp Ser Glu Leu Gly Asn Met Asp Pro Ile Val Thr Val
130 135 140

Lys Lys Tyr Phe Gln Gly Ile Tyr Asp Tyr Leu Gln Glu Lys Gly Tyr
145 150 155 160

Ser Asp Cys Ala Trp Glu Ile Val Arg Val Glu Met Met Arg Ala Leu
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180 185 190

Leu Asn Ser Pro
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35 40 45

Gln Leu Cys Leu Lys Asp Arg Ile Asp Phe Lys Phe Pro Trp Lys Arg
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Glu Ser Ile Thr Gln Leu Gln Lys Asp Gln Ala Phe Pro Val Leu Tyr
65 70 75 80

Glu Met Leu Gln Gln Thr Phe Asn Leu Phe His Thr Glu His Ser Ser
85 90 95

Ala Ala Trp Asn Thr Thr Leu Leu Asp Gln Leu Leu Ser Ser Leu Asp
100 105 110

Leu Gly Leu Arg Arg Leu Glu His Met Lys Lys Asp Asn Met Asp Cys
115 120 125

Pro His Val Gly Ser Ala Leu Arg Lys Tyr Phe Gln Gly Ile Gly Leu
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Val Glu Ile Glu Arg Cys Phe Ser Leu Thr
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35 40 45

Ser Ser Glu Arg Ile Asp Lys Gln Ile Arg Tyr Ile Leu Asp Gly Ile
50 55 60

Ser Ala Leu Arg Lys Glu Thr Cys Asn Lys Ser Asn Met Cys Glu Ser
65 70 75 80

Ser Lys Glu Ala Leu Ala Glu Asn Asn Leu Asn Leu Pro Lys Met Ala
85 90 95

Lys Glu Asp Gly Cys Phe Gln Ser Gly Phe Asn Glu Glu Thr Cys Leu
100 105 110

Val Lys Ile Ile Thr Gly Leu Leu Glu Phe Glu Val Tyr Leu Glu Tyr
115 120 125

Leu Gln Asn Arg Phe Glu Ser Ser Glu Glu Gln Ala Arg Ala Val Gln
130 135 140

Met Ser Thr Lys Val Leu Ile Gln Phe Leu Gln Lys Lys Ala Lys Asn
145 150 155 160

Leu Asp Ala Ile Thr Thr Pro Asp Pro Thr Thr Asn Ala Ser Leu Leu
165 170 175

Thr Lys Leu Gln Ala Gln Asn Gln Trp Leu Gln Asp Met Thr Thr His
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Leu Arg Gln Met
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Arg Ser Leu Ala Leu Cys Gln Lys Leu Leu Arg Gln Leu Pro Ser Thr
35 40 45

Pro Gln His Cys Leu Glu Ala Arg Met Asp Phe Gln Met Pro Glu Glu
50 55 60

Met Lys Gln Ala Gln Gln Phe Gln Lys Glu Asp Ala Ile Leu Val Ile
65 70 75 80

Tyr Glu Met Leu Gln Gln Ile Phe Asn Ile Leu Thr Arg Asp Phe Ser
85 90 95

Ser Thr Gly Trp Ser Glu Thr Ile Ile Glu Asp Leu Leu Glu Glu Leu
100 105 110

Tyr Glu Gln Met Asn His Leu Glu Pro Ile Gln Lys Glu Ile Met Gln
115 120 125

Lys Gln Asn Ser Thr Met Gly Asp Thr Thr Val Leu His Leu Arg Lys
130 135 140

Tyr Tyr Phe Asn Leu Val Gln Tyr Leu Lys Ser Lys Glu Tyr Asn Arg
145 150 155 160

Cys Ala Trp Thr Val Val Arg Val Gln Ile Leu Arg Asn Phe Ser Phe
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Leu Thr Arg Leu Thr Gly Tyr Leu Arg Glu
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Gln Ser Leu Lys Glu Cys Gln Lys Leu Leu Gly Gln Leu Pro Ser Thr
35 40 45

Ser Gln His Cys Leu Glu Ala Arg Met Asp Phe Gln Met Pro Glu Glu
50 55 60

Met Lys Gln Glu Gln Gln Phe Gln Lys Glu Asp Ala Ile Leu Val Met
65 70 75 80

Tyr Glu Val Leu Gln His Ile Phe Gly Ile Leu Thr Arg Asp Phe Ser

                    85   90   95

Ser Thr Gly Trp Asn Ser Thr Thr Glu Asp Thr Ile Val Pro His Leu
            100   105   110

Gly Lys Tyr Tyr Phe Asn Leu Met Gln Tyr Leu Glu Ser Lys Glu Tyr
        115   120   125

Asp Arg Cys Ala Trp Thr Val Val Gln Val Gln Ile Leu Thr Asn Val
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            20   25   30

Lys Asp Arg Met Asn Phe Asp Ile Pro Glu Glu Ile Lys Gln Leu Gln
        35   40   45

Gln Phe Gln Lys Glu Asp Ala Ala Leu Thr Ile Tyr Glu Met Leu Gln
    50   55   60

Asn Ile Phe Ala Ile Phe Arg Gln Asp Ser Ser Ser Thr Gly Trp Asn
65   70   75   80

Glu Thr Ile Val Glu Asn Leu Leu Ala Asn Val Tyr His Gln Ile Asn
                85   90   95

His Leu Lys Thr Val Leu Glu Glu Lys Leu Glu Lys Glu Asp Phe Thr
            100   105   110

Arg Gly Lys Leu Met Ser Ser Leu His Leu Lys Arg Tyr Tyr Gly Arg
        115   120   125

Ile Leu His Tyr Leu Lys Ala Lys Glu Tyr Ser His Cys Ala Trp Thr
    130   135   140

Ile Val Arg Val Glu Ile Leu Arg Asn Phe Tyr Phe Ile Asn Arg Leu
145   150   155   160

Thr Gly Tyr Leu Gly Asn
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ctgccgcaag aattcctgca atacacccag ccgatgaaac gtgatatcaa gaaagcgttc 180
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